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OM nucleic - nucleic search, using sw model

Run on: January 9, 2002, 20:59:50 ; Search time 81.87 Seconds
(without alignments)
10199.394 Million cell updates/sec

Title: US-09-117-447-1
Perfect score: 3687
Sequence: 1 atgatatgaaaaaacgtgt.....ttacattctataataagtaa 3687

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 351203 seqs, 113238999 residues

Total number of hits satisfying chosen parameters: 702406

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Issued_Patents_NA:*

- 1: /cgn2_6/ptodata/2/ina/5A_COMB.seq:*
- 2: /cgn2_6/ptodata/2/ina/5B_COMB.seq:*
- 3: /cgn2_6/ptodata/2/ina/6A_COMB.seq:*
- 4: /cgn2_6/ptodata/2/ina/6B_COMB.seq:*
- 5: /cgn2_6/ptodata/2/ina/PCTUS_COMB.seq:*
- 6: /cgn2_6/ptodata/2/ina/backfiles1.seq:*

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Query Length	DB ID	Description
1	56.8	1.5	4248	3	US-08-678-614-1 Sequence 1, Appli
2	56.2	1.5	397	3	US-09-253-691-3 Sequence 3, Appli
3	53.4	1.4	7218	1	US-08-232-463-14 Sequence 14, Appli
4	51.4	1.4	7291	4	US-08-913-942-3 Sequence 3, Appli
5	51	1.4	2241	2	US-08-785-431-3 Sequence 3, Appli
6	51	1.4	2241	4	US-09-205-048-3 Sequence 3, Appli
7	51	1.4	2367	2	US-08-785-431-1 Sequence 1, Appli
8	51	1.4	2367	4	US-09-205-048-1 Sequence 1, Appli
9	49.8	1.4	5738	1	US-08-409-995-3 Sequence 3, Appli
10	49.8	1.4	5738	3	US-08-685-467-3 Sequence 3, Appli
11	49	1.3	5361	4	US-08-973-462-2 Sequence 2, Appli
12	49	1.3	6152	4	US-08-973-462-1 Sequence 1, Appli
13	48.8	1.3	2223	1	US-08-257-073-4 Sequence 4, Appli
14	47.4	1.3	3095	6	5231168-1 Patent No. 5231168
15	46.6	1.3	4766	5	PCT-US93-07261-10 Sequence 10, Appli
16	45.2	1.2	921	2	US-08-795-475-2 Sequence 2, Appli
17	45.2	1.2	1308	2	US-08-795-475-4 Sequence 4, Appli
18	45.2	1.2	5511	3	US-08-928-361B-2 Sequence 2, Appli
19	45.2	1.2	7334	3	US-08-928-361B-1 Sequence 1, Appli
20	45	1.2	3279	4	US-08-446-137B-1 Sequence 1, Appli
21	44.6	1.2	14602	1	US-08-597-236-1 Sequence 1, Appli
22	44.6	1.2	14602	1	US-08-746-682A-1 Sequence 1, Appli
23	44.4	1.2	1867	1	US-07-772-087-1 Sequence 1, Appli
24	43.8	1.2	1891	4	US-08-973-462-3 Sequence 3, Appli
25	43.6	1.2	1555	3	US-08-669-408B-9 Sequence 9, Appli
26	43.2	1.2	876	4	US-08-446-137B-3 Sequence 3, Appli
27	43.2	1.2	4197	2	US-08-682-517-7 Sequence 7, Appli

28	43.2	1.2	4197	2	US-08-682-517-8	Sequence 8, Appli
29	43.2	1.2	19124	2	US-08-487-826B-13	Sequence 13, Appli
30	42.6	1.2	835	3	US-08-961-083-93	Sequence 93, Appli
31	42.6	1.2	1714	3	US-08-961-083-3	Sequence 3, Appli
32	42.4	1.1	372	1	US-08-094-128A-24	Sequence 24, Appli
33	42.4	1.1	372	1	US-08-455-674-24	Sequence 24, Appli
34	42.4	1.1	372	1	US-08-455-992-24	Sequence 24, Appli
35	42.4	1.1	372	1	US-08-455-972-24	Sequence 24, Appli
36	42.4	1.1	372	5	PCT-US92-00652-24	Sequence 24, Appli
37	42.4	1.1	4977	1	US-08-450-257-14	Sequence 14, Appli
38	42.4	1.1	4977	1	US-08-450-246-14	Sequence 14, Appli
39	42.4	1.1	4977	1	US-08-450-098-14	Sequence 14, Appli
40	42.4	1.1	4977	1	US-08-451-233-14	Sequence 14, Appli
41	42.4	1.1	4977	1	US-08-450-236-14	Sequence 14, Appli
42	42.4	1.1	7904	1	US-08-316-239B-1	Sequence 1, Appli
43	42.4	1.1	7904	1	US-08-316-239B-2	Sequence 2, Appli
44	42.4	1.1	7904	1	US-08-410-005-1	Sequence 1, Appli
45	42.4	1.1	7904	3	US-08-929-140-1	Sequence 1, Appli

ALIGNMENTS

RESULT 1
US-08-678-614-1
: Sequence 1, Application US/08678614
: Patent No. 6013507
: GENERAL INFORMATION:
: APPLICANT: Tomasz, Alexander
: APPLICANT: Delencastre, Herminia
: TITLE OF INVENTION: AUXILIARY GENES AND PROTEINS OF
: NUMBER OF SEQUENCES: 1
: CORRESPONDENCE ADDRESSES:
: ADDRESSEE: David A. Jackson, Esq.
: STREET: 411 Hackensack Ave, Continental Plaza, 4th
: CITY: Floor
: STATE: Hackensack
: STATE: New Jersey
: COUNTRY: USA
: ZIP: 07601
: COMPUTER READABLE FORM:
: MEDIUM TYPE: Floppy disk
: COMPUTER: IBM PC compatible
: OPERATING SYSTEM: PC-DOS/MS-DOS
: SOFTWARE: PatentIn Release #1.0, Version #1.30
: CURRENT APPLICATION DATA:
: APPLICATION NUMBER: US/08/678,614
: FILING DATE: 10-JUL-1996
: CLASSIFICATION: 514
: ATTORNEY/AGENT INFORMATION:
: NAME: Jackson Esq., David A.
: REGISTRATION NUMBER: 26,742
: REFERENCE/DOCKET NUMBER: 600-1-139
: TELECOMMUNICATION INFORMATION:
: TELEPHONE: 201-487-5800
: TELEFAX: 201-343-1684
: INFORMATION FOR SEQ ID NO: 1:
: SEQUENCE CHARACTERISTICS:
: LENGTH: 4248 base pairs
: TYPE: nucleic acid
: STRANDEDNESS: double
: TOPOLOGY: linear
: MOLECULE TYPE: DNA (genomic)
: HYPOTHETICAL: NO
: ANTI-SENSE: NO
: ORIGINAL SOURCE:
: ORGANISM: Staphylococcus aureus
: STRAIN: RUSA 266
: FEATURE:
: NAME/KEY: ORF
: LOCATION: 154..1410
: FEATURE:

Matches 169; Conservative 0; Mismatches 188; Indels 0; Gaps 0;

OY	1454	cagcaccacactgltgccllacccgtccctactactcttagtgatgaacaaccttatactactggtc	1511
Dd	358	CCGCTGGTGCCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG	299
OY	1514	ctcttacacaacaatgltttgggggtaaatgttcgtggtggtgaaatgaagctygaactatt	1573
Dd	298	CTGCTGTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG	239
OY	1574	atcctgytcttcaatlccaacaacaacylltgctaactaagttagaacgaatcactltagctg	1633
Dd	238	CTGCTGTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG	179
OY	1634	ataacttgttatlagttgaaaaagaatatctggtacagtlgttgcttcctgaactaaaatala	1693
Dd	178	CTGCTGTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG	119
OY	1694	atgcagacgctaaaatgyfaacttttagtgccaaaagcggaaccttaagaaatatcaatct	1753
Dd	118	TCTGAACATTCAAAAGTAGAATAATTAAAAAAACAANAACTTAAAGAATAAATACACC	59
OY	1754	atcaaatcaaaatttaaaaaggcttgaagtcgcgataaagyttattgaattagycactg	1810
Dd	58	ATGAGAAAAACTATTCTCATTAAGGAAAATACATGTTTCAACGAAATCAAAGTAGTCACTG	2

RESULT 3

```

US-08-232-463-14/c
; Sequence 14, Application US/08232463
; Patent No. 5670367
; GENERAL INFORMATION:
; APPLICANT: DORNER, F.
; APPLICANT: SCHEIFLINGER, F.
; APPLICANT: FALKNER, F. G.
; TITLE OF INVENTION: RECOMBINANT FOWLPOX VIRUS
; NUMBER OF SEQUENCES: 52
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Foley & Lardner
; STREET: 1800 Diagonal Road, Suite 500
; CITY: Alexandria
; STATE: VA
; COUNTRY: USA
; ZIP: 22313-0299
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/232,463
; FILING DATE:
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US/07/935,313
; FILING DATE:
; APPLICATION NUMBER: EP 91 114 300.6
; FILING DATE: 26-AUG-1991
; ATTORNEY/AGENT INFORMATION:
; NAME: BENT, Stephen A.
; REGISTRATION NUMBER: 29,768
; REFERENCE/DOCKET NUMBER: 30472/114 IMMU
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (703)836-9300
; TELEFAX: (703)683-4109
; TELEX: 899149
; INFORMATION FOR SEQ ID NO: 14:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 7218 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; IMMEDIATE SOURCE:

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CLONE: PTZgpt-F1S
US-08-232-463-14

Query Match 1.4%; Score 53.4; DB 1; Length 7218;
Best Local Similarity 6.3%; Pred. No. 0.0027;
Matches 27; Conservative 222; Mismatches 178; Indels 0; Gaps 0

OY	97	gatgtagcaacagttagtaaggccaagaacaatgcatactactaac	156
Db	1439	GGTACRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRRR ::: :::	1380
OY	157	agccatacagtaacygaaaactygltaatlccccaaacattaacgatatgtctaatac	216
Db	1379	RRR ::: :::	1320
OY	217	aacaaagcgaaaaaaacgaalaccgtlgtcgtltagcatlgtlaataaagcagtgccg	276
Db	1319	RRR ::: :::	1260
OY	277	aaaaagacgcttacctagctgtattacaanaagatatgaacttacgtttcbaagca	336
Db	1259	RR ::: :::	1200
OY	337	aaccctaactctgcgcgaagctcgtlgtlagcaactlacatcgalgtcttacactatgccaa	396
Db	1199	RR ::: :::	1140
OY	397	aathtagcgaatatgcgccaaagctagagcctgctgtltcaaagcaaaagatttagaaaaa	456
Db	1139	RR ::: :::	1080
OY	457	gcagaacaatactatcacaaaaattcctlatgaaattaaactcgcacagtcattttaagat	516
Db	1079	RRRRRRRRRRRRRRATGCCAAGCTCCCTGCACCTGCAACCACAAGCTCGGAATTAAITCIGTG ::: :::	1020
OY	517	cgcgttat 523	
Db	1019	AGCGTAT 1013	

RESULT 4

US-08-913-942-3
: Sequence 3, Application US/08913942
: Patent No. 6200578
: GENERAL INFORMATION:
: APPLICANT: St. Geme, Joseph
: APPLICANT: Barenkamp, Stephen J.
: TITLE OF INVENTION: HAEMOPHILUS ADHESION PROTEINS
: NUMBER OF SEQUENCES: 19
: CORRESPONDENCE ADDRESS:
: ADDRESSEE: Flehr Hohbach Test Albritton & Herbert LLP
: STREET: Four Embarcadero Center, Suite 3400
: CITY: San Francisco
: STATE: California
: COUNTRY: United States
: ZIP: 94111-4187
: COMPUTER READABLE FORM:
: MEDIUM TYPE: Floppy disk
: COMPUTER: IBM PC compatible
: OPERATING SYSTEM: PC-DOS/MS-DOS
: SOFTWARE: PatentIn Release #1.0, Version #1.30
: CURRENT APPLICATION DATA:
: APPLICATION NUMBER: US/08/913,942
: FILING DATE: 29-DEC-1997
: CLASSIFICATION: 514
: PRIOR APPLICATION DATA:
: APPLICATION NUMBER: US 08/409,995
: FILING DATE: 24-MAR-1995
: PRIOR APPLICATION DATA:
: APPLICATION NUMBER: PCT/US96/4031
: FILING DATE: 22-MAR-1996
: ATTORNEY/AGENT INFORMATION:
:

NAME: Vance, Dolly A.
REGISTRATION NUMBER: 39,054
REFERENCE/DOCKET NUMBER: A-61053-1/REF/RMS/DAV
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
TELEFAX: (415) 398-3249
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 7291 base pairs
TYPE: nucleic acid
STRANDEDNESS: unknown
TOPOLOGY: unknown
MOLECULE TYPE: DNA
FEATURE:
NAME/KEY: CDS
LOCATION: 163..7221
US-08-913-942-3

Query Match 1.4%; Score 51.4; DB 4; Length 7291;
Best Local Similarity 45.3%; Pred. No. 0.008;
Matches 276; Conservative 0; Mismatches 321; Indels 12; Gaps 2;

QY 601 gatattaccgttgcaatgaagaagcgcgcaagtagacaagcgtgtgaaagcagcaattta 660
DB 1714 GGTACGACAGGTACAAACCATCAGCGTAACCAAGATGGCATTAAAGCAGGTAATAA 1773
QY 661 gacaaagctaaagctgtgtgtatcaaatcaatcaatcaatcaatcaatcaatcaat 720
DB 1774 GCTATTACTAATGTGGCAGCTGTTAAGAGCTTATGACGATGCGAATTT--TGATGTT 1830
QY 721 ttcaaaactgaactaagaagtagcgaaaaaagcattagatgcagatgaagctgcgctt 780
DB 1831 TTTAATTAACCTGCAACTGATTTAATATGACACGTTGAAGATGCTTATAAAGTTTATTA 1890
QY 781 actccaaaagltgaagtgtaagtgcgalttaacactcaaaaacaaagctgttgaatlaa 840
DB 1891 AATCTAATGAAAAAATGCAAAAT-----AAACAACCGTTGGTGACTGACAGCAGC 1941
QY 841 gcagtaaccagtgaacggaacactaaatlacacattcagctgtgcaaatgaagatata 900
DB 1942 GCGCGGACTGTAGCGGATTTACGTAATTTGGGTTGGGTAGTATCAACCAAAAACGGTACG 2001
QY 901 gtaaacgtaaatcgtacgtatctataaagtgtgacgtgaacattccattgcccattat 960
DB 2002 AAAGAAGAAAGCAATCAAGTTAAACAAGCTGATGAGTCTCTTTACCGGAGCCGGTCT 2061
QY 961 accgcagatgttcttctatctacagacggaacaaactcaactgtgcatgttcaactcca 1020
DB 2062 GCTACGGTTACTTCCAAATCTGAAAAACGGTAACATACGATTACCGTTAGTGGGTGAA 2121
QY 1021 ttcgaaaataatagagataaagtagtagttaaaggtatlaaagacaaaatgycaaa 1080
DB 2122 ACTAAAGCGGATGCGGTCTTGAAAAAGATGGCGATTAATTAAGCTCAAAAGTGGAATAT 2181
QY 1081 gaatttaagaagatgcatcttcaacttcaagcttcgnaatgatgtgtgtactcaagt 1140
DB 2182 CAAAACACTGATTAATGTTTAACTGTTGTAATTAATGTAATGTAATGTAATGTAATG 2241
QY 1141 ttggaactaatgtlaacaacaacacacttctgttaacttagcagcaggtacttgcacact 1200
DB 2242 TTGGAACCTGTTAAACCTGGAAGCAGCTGATGACATCGCGGTAAGTAATGTAATGAT 2301
QY 1201 gacgatact 1209
DB 2302 GCTACTGCT 2310

RESULT 5
US-08-785-431-3
; Sequence 3, Application US/08785431
; Patent No. 5891667

GENERAL INFORMATION:
APPLICANT: Chalke, Alison
APPLICANT: Hodgson, John
TITLE OF INVENTION: No. 5891667el Spoille
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline Beecham Corporation
STREET: 709 Swedeland Road
CITY: King of Prussia
STATE: PA
COUNTRY: USA
ZIP: 19406-0939
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSeq for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/785,431
FILING DATE: 17-JAN-1997
CLASSIFICATION: 530
PRIORITY APPLICATION DATA:
APPLICATION NUMBER: 9600955.0
FILING DATE: 17-JAN-1996
ATTORNEY/AGENT INFORMATION:
NAME: Gimm, Edward R
REGISTRATION NUMBER: 38,891
REFERENCE/DOCKET NUMBER: P31349
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-270-4478
TELEFAX: 610-270-5090
TELEX:
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 2241 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
MOLECULE TYPE: Genomic DNA
US-08-785-431-3

Query Match 1.4%; Score 51; DB 2; Length 2241;
Best Local Similarity 47.8%; Pred. No. 0.0063;
Matches 181; Conservative 0; Mismatches 195; Indels 3; Gaps 1;

QY 2195 atgcaaatcacttagcaactaatatatactatacattcaaacactgaaggtcaagcgttaa 2254
DB 701 ATCGAGAGATAGTTCGAATTAACATCGTAATCATCAAGCAGATCAGCAAGACGAATTA 760
QY 2255 cagcaccacaggttacaaagiatlcaaaagtgatctttaaagacgctgatgcagttta 2314
DB 761 CAGAACAAACTCATACAGTCTTGAAAGTGAAAAACACTATTTGAAGAAGCTGTGAGTTA 820
QY 2315 ctacactlacyaagctgtgacaggtlcaaaaattcactatccaatttagcgaagaattaa 2374
DB 821 CGAATGTATCGTATGTGTCCACCGTTAACTTTACTTTAATCAACCTGCAAAACAAAAG 880
QY 2375 aaacttctagtgttctttagtgggtgcaagtaactgtcgagaatttaacaacaacag 2434
DB 881 CAACATCTTAAGCTGAAGTCAACGTTAAAGGACACAGTACAGATACATTAAGATTT 940
QY 2435 gatgggtagatgctgttactgtgaacaactgtatcagttgtctcctaagacagatgcaaat 2494
DB 941 TTGGGGTAAATGCA--AAAGTACACAAATTAATTTGGTCTCGAGTAACATCAATATG 997
QY 2495 gtaaaaglaacagctgtgtgtltaacatttaactgtgtgacaataaagacaagaatgcga 2554
DB 998 AAATTCACACAGCTCAAGGGGTTAAAGTAGAGTAATAAATTTGTAACCTTGCAATATGAT 1057
QY 2555 aattgcgtctgttagtaga 2573
DB 1058 CATTAAGCTTTAGCAGCAAA 1076


```
RESULT 6
US-09-205-048-3
; Sequence 3, Application US/09205048
; Patent No. 6258936
; GENERAL INFORMATION:
; APPLICANT: Hodgson, John E.
; APPLICANT: Lawlor, Elizabeth J.
; TITLE OF INVENTION: NO. 6258936el spoIIIE
; FILE REFERENCE: P31349-D1
; CURRENT APPLICATION NUMBER: US/09/205,048
; CURRENT FILING DATE: 1998-12-04
; EARLIER APPLICATION NUMBER: 08/785,431
; EARLIER FILING DATE: 1997-01-17
; NUMBER OF SEQ ID NOS: 4
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 3
; LENGTH: 2241
; TYPE: DNA
; ORGANISM: Staphylococcus aureus
US-09-205-048-3
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Query Match 1.4%; Score 51; DB 4; Length 2241;
Best Local Similarity 47.8%; Pred. No. 0.0063;
Matches 181; Conservative 0; Mismatches 195; Indels 3; Gaps 1;

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OY 2195 atgcaaatcttagcaactatatactatatacttaactgaagtcgaagctaa 2254
      ||||| | | | | | | | | | | | | | | | | | | | | | | | | | |
DB 701 atgcagagaatagttcgaataacatcgttaaatcatcagcagatcagcaagcaattaa 760
OY 2255 cagcaccacacggttacaaaagtattcaaaagtgattctttaaaagacgctgatgcgtta 2314
      ||| | | | | | | | | | | | | | | | | | | | | | | | | | |
DB 761 cagaacaactcatacagcgttgaaagtgaaaacacattgaagaagctgtgaaagta 820
OY 2315 ctacaactacgaacgttgatgcaggttcaaaaattcactatccaatttagcgaagaattaa 2374
      | | | | | | | | | | | | | | | | | | | | | | | | | | | |
DB 821 cgaatgtatcgtatgtgtccaccgcttaacttacttaactcaacctgcacaaacaaaag 880
OY 2375 aaacttctagtgttctcttagtgggtgcaaaagtaactgtcgaagaatttaacaacaacg 2434
      ||| |||| | | | | | | | | | | | | | | | | | | | | | |
DB 881 caacatctaagctgaagtaacaacgttaaagacacagtaactagagaatatacattaaagatt 940
OY 2435 gatgggtagatgtgtgactggaacaactgtatcagttgtctcctaagaacagatgcgaatg 2494
      ||||| |||| | | | | | | | | | | | | | | | | | | | | |
DB 941 ttggggtaatatgca--aaagtgcacacaaattaaatgtgtcctgcagtaactcaatatg 997
OY 2495 gtaaaagtaacagctgtgtgtgttacattaaactgtgtcttgacaataaagcacaagaatgca 2554
      | | | | | | | | | | | | | | | | | | | | | | | | | | |
DB 998 aaattcaaccagctcaaggggttaaaagtgaagtaaaaattgtaaaacttgacataatgatattg 1057
OY 2555 aattgcgtctgtgtagtaga 2573
      ||| | | | | | | | |
DB 1058 cattagcttagcagcaaa 1076
```

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RESULT 7
US-08-785-431-1
; Sequence 1, Application US/08785431
; Patent No. 5891667
; GENERAL INFORMATION:
; APPLICANT: Chalker, Allison
; APPLICANT: Hodgson, John
; TITLE OF INVENTION: NO. 5891667el spoIIIE
; NUMBER OF SEQUENCES: 4
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: SmithKline Beecham Corporation
; STREET: 709 Swedeland Road
; CITY: King of Prussia
; STATE: PA
; COUNTRY: USA
; ZIP: 19406-0939
```

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; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
; SOFTWARE: FastSeq for Windows Version 2.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/785,431
; FILING DATE: 17-JAN-1997
; CLASSIFICATION: 530
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 9600955.0
; FILING DATE: 17-JAN-1996
; ATTORNEY/AGENT INFORMATION:
; NAME: Gimm, Edward R
; REGISTRATION NUMBER: 38,891
; REFERENCE/DOCKET NUMBER: P31349
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 610-270-4478
; TELEFAX: 610-270-5090
; TELEX:
; INFORMATION FOR SEQ ID NO: 1:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 2367 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: double
; TOPOLOGY: linear
; MOLECULE TYPE: Genomic DNA
US-08-785-431-1
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Query Match 1.4%; Score 51; DB 2; Length 2367;
Best Local Similarity 47.8%; Pred. No. 0.0064;
Matches 181; Conservative 0; Mismatches 195; Indels 3; Gaps 1;

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OY 2195 atgcaaatcttagcaactatatactatatacttaactgaagtcgaagctaa 2254
      ||||| | | | | | | | | | | | | | | | | | | | | | | | | | |
DB 827 atgcagagaatagttcgaatatacgttaaatcatcagcagatcagcaagcaattaa 886
OY 2255 cagcaccacacggttacaaaagtattcaaaagtgattctttaaaagacgctgatgcgtta 2314
      ||| | | | | | | | | | | | | | | | | | | | | | | | | | |
DB 887 cagacacaaactcatatacagctgttgaaggtaaaacacatttgaaagaagctgtgaagttta 946
OY 2315 ctacaactacgaacgttgatgcaggttcaaaaattcactatccaatttagcgaagaattaa 2374
      | | | | | | | | | | | | | | | | | | | | | | | | | | | |
DB 947 cgatgtatcgtatgtgtccaccgcttaactttactttaattcaacctgcacaaacaaaag 1006
OY 2375 aaacttctagtgttctcttagtgggtgcaaaagtaactgtcgaagaatttaacaacaacg 2434
      ||| |||| | | | | | | | | | | | | | | | | | | | | | |
DB 1007 caacatctaagctgaagtaacaacgttaaagacacagtaactagagaatatacattaaagatt 1066
OY 2435 gatgggtagatgtgtgactggaacaactgtatcagttgtctcctaagaacagatgcgaatg 2494
      ||||| |||| | | | | | | | | | | | | | | | | | | | | |
DB 1067 ttggggttaaatgca--aaagtgcacacaaattaaatgtgtcctgcagtaactcaatatg 1123
OY 2495 gtaaaagtaacagctgtgtgtgttacattaaactgtgtcttgacaataaagcacaagaatgca 2554
      | | | | | | | | | | | | | | | | | | | | | | | | | | |
DB 1124 aaattcaaccagctcaaggggttaaaagtgaagtaaaaattgtaaaacttgacataatgatattg 1183
OY 2555 aattgcgtctgtgtagtaga 2573
      ||| | | | | | | | |
DB 1184 cattagcttttagcagcaaa 1202
```

```
RESULT 8
US-09-205-048-1
; Sequence 1, Application US/09205048
; Patent No. 6258936
; GENERAL INFORMATION:
; APPLICANT: Hodgson, John E.
; APPLICANT: Lawlor, Elizabeth J.
; TITLE OF INVENTION: NO. 6258936el spoIIIE
; FILE REFERENCE: P31349-D1
; CURRENT APPLICATION NUMBER: US/09/205,048
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; CURRENT FILING DATE: 1998-12-04
; EARLIER APPLICATION NUMBER: 08/785,431
; EARLIER FILING DATE: 1997-01-17
; NUMBER OF SEQ ID NOS: 4
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 1
; LENGTH: 2367
; TYPE: DNA
; ORGANISM: Staphylococcus aureus
;
US-09-205-048-1

```

Query Match	1.4%;	Score 51;	DB 4;	Length 2367;
Best Local Similarity	47.8%;	Pred. No. 0.0064;		
Matches 181; Conservative	0;	Mismatches 195;	Indels 3;	Gaps 1;

QY	2195	atgcaaatactttgagcaactaaactatatactatatacttaacaaactgaaaggtcaagacgttaa	2254
Db	827	atgcagagaaatagtttcgataataacatcgttaaatcatcaagcagatcagcaagagcaattaa	886
QY	2255	cagcaccacaacggttaacaaaaglatccaaggtgatlcttlaaaagacgctgatagcagtta	2314
Db	887	cagaacaactcatcaacagtgltgnaagtgnaaacacatttgaagaagctggtgaagtta	946
QY	2315	ctaacacttacgaacgttgcataagtcaggtcacaanaattcaactccaatttagcgaagaattaa	2374
Db	947	cgaatgtatcgtatggtgttcacacggttaactttacttaactcaactgtcaanaacaaaag	1006
QY	2375	aaacttctagtgttctcttagtgyggtgycaaaglaacgttcgagaataataacaacaacg	2434
Db	1007	caacatctaagcttgaagatlaacaacgtlaaagacaaagtactagagatacatattaaagatt	1066
QY	2435	gatggtgtagatgctgtgactctggaacaactgtatcagttgcttcctaagacagatgcaaatg	2494
Db	1067	ttgggtgtaaatgca---aaagtgcacacaatlaaataattgtctcctgcagtaactccaatg	1123
QY	2495	gtaaagtaacacagctgctgtgttatacttaactggtccttgacaataacgcacaagaatgcga	2554
Db	1124	aaattcaaccagctccaagggttaaaagtgaagtaaaatctgtaaaactgcataatgtatatg	1183
QY	2555	aattgcgtctggtagtaga	2573
Db	1184	cattagctttagcagcaaa	1202

RESULT 9
 US-08-409-995-3
 ; Sequence 3, Application US/08409995
 ; Patent No. 5646259
 ; GENERAL INFORMATION:
 ; APPLICANT: Barenkamp, Stephen I.
 ; APPLICANT: St. Geme III, Joseph W.
 ; TITLE OF INVENTION: Haemophilus Adhesion Proteins
 ; NUMBER OF SEQUENCES: 6
 ; CORRESPONDENCE ADDRESS:
 ; ADDRESSEE: Flehr, Hohbach, Test, Albritton & Herbert
 ; STREET: Four Embarcadero Center, Suite 3400
 ; CITY: San Francisco
 ; STATE: CA
 ; COUNTRY: USA
 ; ZIP: 94111-4187
 ; COMPUTER READABLE FORM:
 ; MEDIUM TYPE: Floppy disk
 ; COMPUTER: IBM PC compatible
 ; OPERATING SYSTEM: PC-DOS/MS-DOS
 ; SOFTWARE: PatentIn Release #1.0, Version #1.30
 ; CURRENT APPLICATION DATA:
 ; APPLICATION NUMBER: US/08/409,995
 ; FILING DATE: 24-MAR-1995
 ; CLASSIFICATION:
 ; ATTORNEY/AGENT INFORMATION:
 ; NAME: Silva, Robin M.
 ; REGISTRATION NUMBER: 38,304

```

; REFERENCE/DOCKET NUMBER: A-61053/RFT
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (415) 781-1989
; TELEFAX: (415) 398-3249
; TELEEX: 910 277299
; INFORMATION FOR SEQ ID NO: 3:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 5738 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: double
; TOPOLOGY: unknown
;
US-08-409-995-3

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Query Match	1.48;	Score 49.8;	DB 1;	Length 5738;
Best Local Similarity	45.28;	Pred. No. 0.017;		
Matches	275;	Conservative	0;	Mismatches 322;
			Indels	12;
			Gaps	2;

QY	601	galatlaccgttgcgaatgaaagcgcgcggaagtacaaagcgtgtgaaagcaggcaattt	660
Db	1549	GGTAGACAGGTACAAACACCATCAGCGTAAACCAAGATGGCATTAAAGCAGGTAATAA	1608
QY	661	gacaaagctaaagctgctgttgcataatcaatcaatcttaccaaaagtaacagatgct	720
Db	1609	GCTATTACTAATGTGGCAGTGGGTTAAGAGCGTTATGACGATGGCAATTTT--TGAATGTT	1665
QY	721	ttcaaaactgaactaacagaagtagcgaaaaaagcattagatgcagatgaaagctgcgctt	780
Db	1666	TTAATAACTCTGCACACTGATTTAATATAGACACCGTTGAAGATGCTTATAAAGCTTTATTA	1725
QY	781	actccaaaagltgaaagtgtaaagtgtaagtcgattaaccaactcaaaaacaaagctgttgaatta	840
Db	1726	AATCTAATGAAAAAATGCA-----AATTAACAACCGTTGGTGCATGACACGACG	1776
QY	841	gcagtaaccagtgaacggaacactaaatlacaaclttcagctgtcgcaatgaaatata	900
Db	1777	GCGGCGACTGTAGGCGATTTACGTAAATTGGGTTGGGTAGTATCAACCAAAAAACGGTTACG	1836
QY	901	gtaaacgtlaaatactgttacgtatctataaaagltgacggttaacalttccatttgccctta	960
Db	1837	AAAGAGAAGAACCAATCAGTTTAACAACAGCTGATGAGTTCCTTTACCGGAGCCGGTTCCT	1896
QY	961	acggcagatgttcttctatctacagacgcgaaaaactatactgctgtgatgcttcaactcca	1020
Db	1897	GCTACGGTTACTTCCAAATCTGAAAAACGGTAAACATACGATTACCGTTAGTGTGGCTGAA	1956
QY	1021	ttcgaaaataatcagaglataaagtagttagttaaagttataaagcaaaaatgycaaa	1080
Db	1957	ACTAAAGCGGATTCGGCTCTTGAAAAAGATGGCGATCTACTATTAAAGCTCAAAAGTGGATTAAT	2016
QY	1081	gaatttaaagaagatgcatttcaacttccaagcttcgaatgatgctgttagtctactcaagtg	1140
Db	2017	CAAAACACTGATTAATGTTTAACTGTTGGTAATTAATGCTACTGTCACATAAAGTGCGC	2076
QY	1141	tttggaaactaatgtaacacaacaacacttctgtaaacttagcagcaggtactcttgcacact	1200
Db	2077	TTTGAAACTGTTAAAACTGCAAGCGCACTGATGCAGATCCCGCTAAAGTAACGTGTAAGAAT	2136
QY	1201	gacgatact 1209	
Db	2137	GCTACTGCT 2145	

RESULT 10
 US-08-685-467-3
 ; Sequence 3, Application US/08685467
 ; Patent No. 6060059
 ; GENERAL INFORMATION:
 ; APPLICANT: St. Geme III, Joseph W.
 ; APPLICANT: Barenkamp, Stephen J.
 ; TITLE OF INVENTION: HAEMOPHILUS ADHESION PROTEINS
 ; NUMBER OF SEQUENCES: 6
 ; CORRESPONDENCE ADDRESS:
 ;
 ;

```

; ADDRESS: Flehr, Hohbach, Test, Albritton & Herbert
; STREET: Four Embarcadero Center, Suite 3400
; CITY: San Francisco
; STATE: California
; COUNTRY: United States
; ZIP: 94111-4187
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/685,467
; FILING DATE: 22-JUL-1996
; CLASSIFICATION: 424
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 08/409,995
; FILING DATE: 24-MAR-1995
; ATTORNEY/AGENT INFORMATION:
; NAME: Silva, Robin M.
; REGISTRATION NUMBER: 38,304
; REFERENCE/DOCKET NUMBER: A-61053-2/RFT/RMS
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (415) 781-1989
; TELEFAX: (415) 398-3249
; TELEX: 910 277299
; INFORMATION FOR SEQ ID NO: 3:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 5738 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: unknown
; TOPOLOGY: unknown
; MOLECULE TYPE: DNA (genomic)
; US-08-685-467-3

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Query Match 1.4%; Score 49.8; DB 3; Length 5738;
Best Local Similarity 45.2%; Pred. No. 0.017;
Matches 275; Conservative 0; Mismatches 322; Indels 12; Gaps 2;

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QY 601 gatattaccgttgcgaagaaagcgcgcgaagtcacgaagcgtgtgaaagcagcgaattta 660
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1549 GGTACGACAGGTACAAACACCATCAGCGTAAACCAAGATGGCATTTAAAGCAGTTAATAA 1608

QY 661 gacaaagctaaagctgctgttgatcaaatcaatcaacttaccaaaagtaacagatgct 720
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1609 GCTATTACTAATGTTGGAGTGGGTTAAGAGCTTATGACGATGCGAATTT--TGATGTT 1665

QY 721 ttcaaaactgaactaacagaagtagcgaaaagaattagatgcagatgaagctgcgctt 780
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1666 TTAATAACTCTGCACTGATTAAATAGACACGTTGAAGATGCTTATTAAGCTTTATTTA 1725

QY 781 actccaaaagltgaagtgtaagtgcgattaacactcaaaaacaaagctgttgaattaca 840
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1726 AATCTAATGAATAAAATGCA-----AATTAACAACCGTTGGTGACTGACACACACG 1776

QY 841 gcagtaaccagtgaacggaacactaaattacaacttcaagctgctgcaaatgaagataca 900
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1777 GCGGCGACTGTAGCGGATTTACGTAATTTGGGTTGGGTAGTATCAACCAAAAACGGTACG 1836

QY 901 gtaacgtaaatactgtacgtatctataaagtgacggtaacattccattgcccctaat 960
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1837 AAAGAAGAAAGCAATCAAGTTAAACAAGCTGATGAAGTCTCTTTACCGGAGCCGGTCT 1896

QY 961 acgycagatgttcttctatctacagacggaanaactatcactgtgcatcttcaactcca 1020
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1897 GCTACGGTTACTTCCAAATCTGAAAAACGGTAAACATACGATTAACCGTTAGTGTGCTGAA 1956

QY 1021 ttcgaaaataatagagataaagtagtagttaaggtalttaagacaaaatgycaaa 1080
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1957 ACTAAAGCGGATTCGGGCTTGAAAAAAGATGGCGATTAATAAGCTCAAAAGTGGATAAT 2016

QY 1081 gaatttaagaagatgcatctcacttccaagcttcgaaatgatgctgtagttactccaagt 1140

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Db 2017 CAAACACTGATATGTTTAACTGTTGGTAATAATGGTACTGCTGCACCTAAAGGTGGC 2076

QY 1141 ttggaaactaatgtacaacaacacttctgtlaaacttagcagcagtagtacttgcacact 1200
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 2077 TTTGAAACTGTTAAAACTGACGCGACTGATGACAGATCGCGTAAAGTAAGTAAAGAT 2136

QY 1201 gacgatact 1209
   ||| |||
Db 2137 GCTACTGCT 2145

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RESULT 11
US-08-973-462-2
; Sequence 2, Application US/08973462B
; Patent No. 6191270
; GENERAL INFORMATION:
; APPLICANT: DRUILHE, PIERRE
; APPLICANT: DAUBERSIES, PIERRE
; TITLE OF INVENTION: MALARIAL PRE-ERYTHROCYTIC STAGE POLYPEPTIDE MOLECULES
; FILE REFERENCE: 0660-0125-0 PCT
; CURRENT APPLICATION NUMBER: US/08/973,462B
; CURRENT FILING DATE: 1998-02-06
; EARLIER APPLICATION NUMBER: PCT/FR96/00894
; EARLIER FILING DATE: 1996-06-12
; EARLIER APPLICATION NUMBER: FR 95/07007
; EARLIER FILING DATE: 1995-06-13
; NUMBER OF SEQ ID NOS: 29
; SOFTWARE: Patent In Ver. 2.0
; SEQ ID NO 2
; LENGTH: 5361
; TYPE: DNA
; ORGANISM: P. falciparum
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (1)..(5361)
; US-08-973-462-2

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Query Match 1.3%; Score 49; DB 4; Length 5361;
Best Local Similarity 45.6%; Pred. No. 0.026;
Matches 215; Conservative 0; Mismatches 250; Indels 6; Gaps 1;

```

QY 1880 aaatgycgacgctgatataaagtaactgaagtcacgaatttactgtgaagttctcag 1939
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1118 aaagtgtagaagaataatgttgaagaagtgtagctgaanaatgttgaagaagtgtagctg 1177

QY 1940 agaatttaatatcatltaatgtctacaaccggttcggttagcacatacatcagtcag 1999
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1178 aaaatgttgaagaagtgtagctgaanaatgttgaagaagtgtagctgaanaatgttgaag 1237

QY 2000 ttgctgtagtaaaagcgggtgcaactatctgctcttaccagcaagtgacatcattccag 2059
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1238 aaagtgtagctgaanaatgttgaagaatactgtagctccaactgttgaagaatcgttagctc 1297

QY 2060 ctagtgttgaagcgttactgtgcaagatggaacatacaaaagt-----aaagttgctg 2113
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1298 caactgttgaagaataatgttagctccaagtgcttgaagaagtgtagctccaagtgtagaag 1357

QY 2114 ctaaccaatttagaacgttaaccaaggtlacaaattagtagtgttcgtaaaagtgcaacag 2173
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1358 aaagtgtagaagaanaatgttgaagaagaagtgtagcttgaanaatgttgaagaagaagtgtagctg 1417

QY 2174 ctccctgttaagaatgtcgtcaaatgtcaaatctttagcaactaactatatactataactt 2233
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1418 aaaatgttgaagaagaagtgtagctgaanaatgttgaagaagaagtgtagctgaanaatgttgaag 1477

QY 2234 caactgaaggtcaagaagtaacagcaacaaaggttacaanaaglatlcaaaagtgatctc 2293
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1478 aaagtgtagctgaanaatgttgaagaagaagtgtagctgaanaatgttgaagaagaagtgtagctg 1537

QY 2294 taaaagcgtgatgcatctactacacttacgaagcgttgatgaggtcaaa 2344
   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||

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Db 1538 aaatgttgaagaatcgtagctcccaactgttgaagaatcgtagctccaa 1588

RESULT 12

US-08-973-462-1

; Sequence 1, Application US/08973462B

; Patent No. 6191270

; GENERAL INFORMATION:

; APPLICANT: DRUILHE, PIERRE

; APPLICANT: DAUBERSIES, PIERRE

; TITLE OF INVENTION: MALARIAL PRE-ERYTHROCYTIC STAGE POLYPEPTIDE MOLECULES

; FILE REFERENCE: 0660-0125-0 PCT

; CURRENT APPLICATION NUMBER: US/08/973,462B

; EARLIER APPLICATION NUMBER: PCT/FR96/00894

; EARLIER FILING DATE: 1996-06-12

; EARLIER APPLICATION NUMBER: FR 95/07007

; EARLIER FILING DATE: 1995-06-13

; NUMBER OF SEQ ID NOS: 29

; SOFTWARE: Patentin Ver. 2.0

; SEQ ID NO 1

; LENGTH: 6152

; TYPE: DNA

; ORGANISM: P. falciparum

US-08-973-462-1

Query Match 1.3%; Score 49; DB 4; Length 6152;

Best Local Similarity 45.6%; Pred. No. 0.027;

Matches 215; Conservative 0; Mismatches 250; Indels 6; Gaps 1;

QY 1880 aaatgcygcgcgtgtgatataaagtgaagctcaagaattactgtgaagttccag 1939

Db 1362 aaagtgtagaagaatgttgaagaagaagtgtagctgtaaaatgttgaagaagaagtgtagctg 1421

QY 1940 agaatttaatacatttaattgtctacaaaccttctcggttagcacaatcacatagctgaag 1999

Db 1422 aaatgttgaagaagaagtgtagctgtaaaatgttgaagaagaagtgtagctgtaaaatgttgaag 1481

QY 2000 ttgctgttagtaaaagcgggtgcaaaacttactgtctcttcaagcaagtgacatcatccag 2059

Db 1482 aaagtgtagctgtaaaatgttgaagaagaatcgtagctccaactgttgaagaagaatcgtagctc 1541

QY 2060 ctagtgttgaagcgttactgtgtcgaagatggaacatcacaaagtg-----aaagtgctg 2113

Db 1542 caactgttgaagaagaatgttagctccaagtggttgaagaagtggtgtccaagtggtgaag 1601

QY 2114 ctaaccaattagaagcgttaaccaaggygtacaatlagtagtggttcgylaaagtgcaacag 2173

Db 1602 aaagtgtagaagaagaatgttgaagaagaagtgtagctgtaaaatgttgaagaagaagtgtagctg 1661

QY 2174 ctctgtttaaagatgtgtcaaatgcaaatactttagcaactaactatatactatatactta 2233

Db 1662 aaatgttgaagaagaagtgtagctgtaaaatgttgaagaagaagtgtagctgtaaaatgttgaag 1721

QY 2234 caactgaagtgcaagcgttaacagcaccaacggttacaaaagtaltcaaaagtgatctt 2293

Db 1722 aaagtgtagctgaagaatgttgaagaagaagtgtagctgtaaaatgttgaagaagaagtgtagctg 1781

QY 2294 taagaagcgtgtagtgcagttactactacactttaggaacggttgatgcaggtcaca 2344

Db 1782 aaatgttgaagaagaatcgttagctccaactgttgaagaagaatcgttagctccaa 1832

RESULT 13

US-08-257-073-4

; Sequence 4, Application US/08257073

; Patent No. 5766597

; GENERAL INFORMATION:

; APPLICANT: Paoletti, Enzo

; APPLICANT: de Taisne, Charles

; APPLICANT: Tine, John A.

; TITLE OF INVENTION: MALARIA RECOMBINANT POXVIRUS VACCINE

; NUMBER OF SEQUENCES: 143

; CORRESPONDENCE ADDRESS:

; ADDRESSEE: Curtis, Morris & Safford, P.C.

; STREET: 530 Fifth Avenue, 25th Floor

; CITY: New York

; STATE: New York

; COUNTRY: UNITED STATES OF AMERICA

; ZIP: 10036

; COMPUTER READABLE FORM:

; MEDIUM TYPE: Floppy disk

; COMPUTER: IBM PC compatible

; OPERATING SYSTEM: PC-DOS/MS-DOS

; SOFTWARE: Patentin Release #1.0, Version #1.30

; CURRENT APPLICATION DATA:

; APPLICATION NUMBER: US/08/257,073

; FILING DATE: 09-JUN-1994

; CLASSIFICATION: 424

; PRIOR APPLICATION DATA:

; APPLICATION NUMBER: US 08/075,783

; FILING DATE: 11-JUN-1993

; PRIOR APPLICATION DATA:

; APPLICATION NUMBER: US 07/852,305

; FILING DATE: 18-MAR-1992

; PRIOR APPLICATION DATA:

; APPLICATION NUMBER: US 07/672,183

; FILING DATE: 20-MAR-1991

; ATTORNEY/AGENT INFORMATION:

; NAME: Frommer, William S.

; REGISTRATION NUMBER: 25,506

; REFERENCE/DOCKET NUMBER: 454310-2570

; TELEPHONE: (212) 840-3333

; TELEFAX: (212) 840-0712

; TELEX: 425066 CURTMS

; INFORMATION FOR SEQ ID NO: 4:

; SEQUENCE CHARACTERISTICS:

; LENGTH: 2223 base pairs

; TYPE: nucleic acid

; STRANDEDNESS: single

; TOPOLOGY: linear

US-08-257-073-4

Query Match 1.3%; Score 48.8; DB 1; Length 2223;

Best Local Similarity 42.1%; Pred. No. 0.021;

Matches 543; Conservative 0; Mismatches 732; Indels 15; Gaps 4;

QY 558 atttaagcaaaagcacaaagacttcgcgaagcttaattatgatattaccgttgcatt 617

Db 534 ATTTAAGAATTATGATGATCAAAAAGCTACCTACTCTTTGTTATATACCAAAAAAATAAT 593

QY 618 gaaagcygcgaagtacaagcgtgtgtgaagcagcgaatttagacaaagctaaagctgc 677

Db 594 TATTTCCTTTAAATCACAAGGTGATTAATAAAGAAACCTCACAAAATCAAAATGAATAA 653

QY 678 tgtgatcaaatcaatcaatctaccacaaagttaacagatgctttcaaaactgaactaac 737

Db 654 CGACAATCAAAAATATATCAAGAAGTTAATGATGAAGATGATGTAATGATGAAGAAGATAC 713

QY 738 agaagtagcgyaaaaaagcatlagatgycagatgaaagctgcgttactccaagaagttgaaag 797

Db 714 AACGATGACGAAGATACTACGATGAAGAAGATACAAACGATGACGAAGATACAAATGA 773

QY 798 tgtaatgcygatttaacactcaaaacaaagcgtgttgaattaaacagcagtlaccagtyaacgy 857

Db 774 TGACGAAGATACTAACGATGAAGAAGATACTAACGACGAAGAAGATCATGAATAATTA 833

QY 858 aacactaaattacaactttagcgtgtgtgcaaatgaaatagatacagtaaacgtaaatcgt 917

Db 834 TCGTACAGCATACGAATTAGTATCTCCAGTTAACGATGTGTAAATGTTAATATGA 893

QY 918 acgtatcataaagtgagcgttaacattccatttgccttaatacgcgcagatgttctt 977


```

Db 894 AAATATGATAACAGGAAA--TAATTTATGATGTTGTTAAAGTACATTAGCTCAATC 950
OY 978 atctacagacggaanaactatcactgtgcttcaactccattcganaataacgga 1037
Db 951 AGGTGATTAGGAATGATGATTTAATTAATTTCTTAATCAAGCTAAAGAAATAGGAGA 1010
OY 1038 gtataagtagtaaggtatataagacaaaatgcaagaattlaaagaagatgc 1097
Db 1011 AAATTTATTAACAATACAAAGATGACCTTGGAGATAGAATAATCTTGAAGCTTTCC 1070
OY 1098 attcaact--ttcaagcttcgaatgatgctgttactcaagtggttgaactaatgt 1154
Db 1071 TTTAGATCAATTAATATGTTAAAGATTAATTAACATGATGATTCATTAATAATAA 1130
OY 1155 aacaacaacactctctgtaaaacttagcagcaggtacttgcacactgcagatacttaac 1214
Db 1131 TTTGAAAACAAGTGTTTAAATTAATTAAGATTTATTTAAGATTATTATACAAAGC 1190
OY 1215 agtagtattgataagttgttagcacctgaaactgtaaacagctcgaaacttactaac 1274
Db 1191 ATATGTATCATATAGAAAAGAAAAGCTCAAGAAAAGGATTACCAAGACCTACTGTTAC 1250
OY 1275 agatgtgaactggaanaacgcatccagtaattgcatctactctgtgttcaactaac 1334
Db 1251 TAATGAGAATATGTTGAA-GAATTAAAGAAAGGTATTCTAGATATGGTATCAAAATTAT 1309
OY 1335 tattacgtttaaagaagcgttagtaactgttaacaatataaactgtctatacataatgt 1394
Db 1310 TATTTAGTAAGTTAAAGCCCTATTAAAAAAATTAATAATTAATTTCC----- 1361
OY 1395 taaaacattactgtgttacaatgcaagaacttaccagttagtggttcaactgcgaacgcatc 1454
Db 1362 TAAGAAAAAGAGAGATTAATCAAGCAGTAGATACCAAAAGTATGGAGAAGACCCAAAGTTAA 1421
OY 1455 agcaccacactgtgtcaccgctcctactactttagtggtgtaacaacttactactgtgtc 1514
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OY 1815 cgagaaacatatgagttcaaaactcaaga 1844
Db 1782 TAAAAAACACATGAGCTATAAACACAGA 1811

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RESULT 14
5231168-1
; Patent No. 5231168
; APPLICANT: DIEGIEL, MORTEN, BORRE, MARTIN, JEPSEN, SOREN;
; VUUST, JENS, RIENECK, KLAUS, WIND, ANNETTE, JAKOBSEN, PALLE H.
; TITLE OF INVENTION: MALARIA ANTIGEN
; NUMBER OF SEQUENCES: 19
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/07/409,658
; FILING DATE: 18-SEP-1989

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; SEQ ID NO:1:
; LENGTH: 3095
5231168-1
Query Match 1.3%; Score 47.4; DB 6; Length 3095;
Best Local Similarity 42.1%; Pred. No. 0.05;
Matches 270; Conservative 0; Mismatches 371; Indels 0; Gaps 0;
OY 417 agagctagagcgtcgttcaagcaaaagalltagaaaaaagcagaacaatactatcacaa 476
Db 1269 agaggttgaagaattctaccagaagaatgatataaaatgaaaaagttgaacatgaataagt 1328
OY 477 aattcctatgaatlaaaactgcacagtcatttagatcgcgtlatatggtlaaacaac 536
Db 1329 agaagltgaagaattctaccagaagataaaaaatgaaaaagltcacatgaataatgtaga 1388
OY 537 tcgtgatttacttcgctctacattttaaagcaaaagcacaagaacttcgcgacagctaat 596
Db 1389 ggttgaagaattctaccagaagataaaaaatgaaaaagltgaacatgaataatgtagaagt 1448
OY 597 ttatgatattaccglttgcaatgaaagcgcggaagtacaaagcgtgtgtaaaagcagcaa 656
Db 1449 tgaagaattctaccagaagataaaaaatgaaaaagltcaacatgaataatgtagaaggttga 1508
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Db 1509 agaattctaccagaagtaaaaaatgaaaaagltcaacatgaataatgtagaaggttgaaga 1568
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Db 1569 aattctaccagaagataaaaaatgaaaaaggtcacaatgaataatgtagaggttgaagaat 1628
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Db 1689 tctaccagaagaataaaaaatgaaaaaggtcacaatgaataatgtagaggttgaagaatctc 1748
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Db 1749 accagaagataaaaaatgaaaaaggtcacaatgaataatgtagaggttgaagaatctc 1808
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Db 1809 agaattgttgaatltgaagaagtagtaccatcacacaacaatacaaatgaaatattgaaac 1868
OY 1017 tccattcgaaaaataatacagaggtataaagtagtagttaag 1057
Db 1869 tataaaaccagaagaataaagaatgaatttagtgttgaag 1909

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RESULT 15
PCT-US93-07261-10
; Sequence 10, Application PC/TUS9307261
; GENERAL INFORMATION:
; TITLE OF INVENTION: PFEHP3 MALARIA ANTIGEN, ANALOGS, ANTIBODIES AND USES THEREO
; NUMBER OF SEQUENCES: 23
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: John H. C. Blasdale
; STREET: One Giralda Farms
; CITY: Madison
; STATE: New Jersey
; COUNTRY: USA
; ZIP: 07940-1000
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: Apple Macintosh
; OPERATING SYSTEM: Macintosh 6.0.5
; SOFTWARE: Microsoft Word 5.1a
; CURRENT APPLICATION DATA:

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APPLICATION NUMBER: PCT/US93/07261
FILING DATE: 19930805
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/927,531
FILING DATE: 07-AUG-1992
ATTORNEY/AGENT INFORMATION:
NAME: Blasdale, John H. C.
REGISTRATION NUMBER: 31,895
REFERENCE/DOCKET NUMBER: DX0288K
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-822-7398
TELEFAX: 201-822-7039
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 4766 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
MOLECULE TYPE: cDNA to mRNA
ORIGINAL SOURCE:
ORGANISM: Plasmodium falciparum
STRAIN: Malayan Camp
IMMEDIATE SOURCE:
CLONE: p2b1;p12-1
FEATURE:
NAME/KEY: CDS
LOCATION: 3..4766
PCT-US93-07261-10
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Query Match 1.38; Score 46.6; DB 5; Length 4766;
Best Local Similarity 42.68; Pred. No. 0.091;
Matches 302; Conservative 0; Mismatches 404; Indels 3; Gaps 1;
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QY 240 tgatgcggtagcatagtgaaataaagcaggtgcggaaaaaagacgcttactagctga 299
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Db 3050 AGGATTAAAGAAAAATGCAGAAATTAAAAAATTAAGAAATTACGAAATTAAGGATCTGAAGG 3109

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Db 3170 AAAAGAAAAATGCAGAACTAAAAAATAAAGAAATTACAAAAATTAAGGATCTGAAGGATTAAA 3229

QY 840 agcagtlaccagtgtaacggaacactaaaattacaactttcagctgctgca 888
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